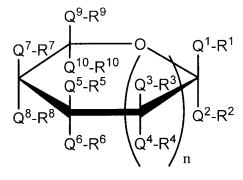
## Amendments to the Claims:

This listing of claims will replace all prior versions, and listings of claims in the application:

## **LISTING OF CLAIMS:**

- 1 1. (Original) A method of forming a peptide conjugate comprising a covalent linkage
- 2 between a modifying group and a glycosylated or non-glycosylated peptide, wherein said
- 3 modifying group is conjugated to the peptide via a glycosyl linking group interposed between
- 4 and covalently linked to both said peptide and said modifying group, said method comprising:
- a. contacting a cell with a modified sugar comprising a sugar moiety and at least
- one modifying group, wherein said modifying group is a member independently selected from
- 7 the group consisting of a water-soluble polymer, a therapeutic moiety, a detectable label, a
- 8 biomolecule and a targeting moiety;
- b. incubating said cell under conditions in which said cell internalizes said
- 10 modified sugar;
- c. after step b, intracellularly contacting said modified sugar with a glycosylated
- or non-glycosylated peptide and a glycosyltransferase for which said modified sugar is a
- substrate, thereby forming said peptide conjugate.
  - 1 2. (Original) The method of claim 1, further comprising, after step b and before step c,
  - 2 intracellularly contacting said modified sugar with a nucleotide and a nucleotidyl transferase,
  - 3 thereby forming a modified nucleotide sugar, wherein
  - 4 said modified sugar in step c is said modified nucleotide sugar.
  - 1 3. (Original) The method of claim 1, further comprising isolating said peptide conjugate.
  - 4. (Original) The method of claim 1, wherein said modified sugar is a modified nucleotide
  - 2 sugar.
  - 1 5. (Original) The method of claim 1, wherein said modified sugar is a modified activated
  - 2 sugar.

- 1 6. (Original) The method of claim 1, wherein said glycosyl linking group is an intact
- 2 glycosyl linking group.
- 1 7. (Original) The method of claim 1, wherein said modified sugar is a precursor modified
- 2 sugar that is intracellularly converted to an intermediate modified sugar by cellular enzymes after
- 3 step b and before step c.
- 8. (Original) The method of claim 7, wherein said intermediate modified sugar is a
- 2 phosphorylated modified sugar, wherein said phosphorylated modified sugar is formed by
- 3 intracellularly contacting said modified sugar with a kinase for which said modified sugar is a
- 4 substrate, thereby forming a phosphorylated modified nucleotide sugar.
- 9. (Original) The method of claim 1, wherein said water-soluble polymer comprises
- 2 poly(ethylene glycol).
- 1 10. (Original) The method of claim 10, wherein said poly(ethylene glycol) has a molecular
- 2 weight distribution that is essentially homodisperse.
- 1 11. (Original) The method of claim 1, wherein said modified sugar has the formula



(I)

wherein,

2

4 n represents an integer from 0 to 1;

Q<sup>1</sup>, Q<sup>2</sup>, Q<sup>3</sup>, Q<sup>4</sup>, Q<sup>5</sup>, Q<sup>6</sup>, Q<sup>7</sup>, Q<sup>8</sup>, Q<sup>9</sup>, and Q<sup>10</sup> are members independently selected

from a bond, substituted or unsubstituted alkylene, substituted or

7 unsubstituted heteroalkylene, substituted or unsubstituted cycloalkylene,

8 substituted or unsubstituted heterocycloalkylene, substituted or unsubstituted

9		arylene, substituted or unsubstituted heteroarylene, -O-, -N(R''')-, -S-, -C(O)-,
10		and -CH <sub>2</sub> -, wherein
11		R <sup>1A</sup> is a member selected from hydrogen, substituted or unsubstituted alkyl,
12		substituted or unsubstituted heteroalkyl, substituted or unsubstituted
13		cycloalkyl, substituted or unsubstituted heterocycloalkyl, substituted or
14		unsubstituted aryl, and substituted or unsubstituted heteroaryl; and
15		R <sup>1</sup> , R <sup>2</sup> , R <sup>3</sup> , R <sup>4</sup> , R <sup>5</sup> , R <sup>6</sup> , R <sup>7</sup> , R <sup>8</sup> , R <sup>9</sup> , and R <sup>10</sup> are members independently selected
16		from -OPO <sub>3</sub> H <sub>2</sub> , -OH, -NH <sub>2</sub> , -SH, hydrogen, substituted or unsubstituted alkyl,
17		substituted or unsubstituted heteroalkyl, substituted or unsubstituted
18		cycloalkyl, substituted or unsubstituted heterocycloalkyl, substituted or
19		unsubstituted aryl, substituted or unsubstituted heteroaryl, an activated leaving
20		group, a nucleotidyl moiety, and a modifying group, wherein at least one of
21		R <sup>1</sup> , R <sup>2</sup> , R <sup>3</sup> , R <sup>4</sup> , R <sup>5</sup> , R <sup>6</sup> , R <sup>7</sup> , R <sup>8</sup> , R <sup>9</sup> , and R <sup>10</sup> is a modifying group.
1	12.	(Original) The method of claim 11, wherein
2		$Q^{1}-R^{1}$ , $Q^{2}-R^{2}$ , $Q^{3}-R^{3}$ , $Q^{4}-R^{4}$ , $Q^{5}-R^{5}$ , $Q^{6}-R^{6}$ , $Q^{7}-R^{7}$ , $Q^{8}-R^{8}$ , $Q^{9}-R^{9}$ , and $Q^{10}-R^{10}$ are
3		members independently selected from hydrogen, -OPO <sub>3</sub> H <sub>2</sub> ,-OH, -OCH <sub>3</sub> , -
4		$CH_3$ , $-C(O)H$ , $-CH_2OH$ , $-NHR^{11}$ , $-O-CH(CH_3)COOR^{12}$ , $-C(O)OR^{13}$ , $-CHR^{14}$ -
5		CHR <sup>15</sup> -CH <sub>2</sub> R <sup>16</sup> , an activated leaving group, a nucleotidyl moiety and -L-M,
6		wherein at least one of R <sup>1</sup> , R <sup>2</sup> , R <sup>3</sup> , R <sup>4</sup> , R <sup>5</sup> , R <sup>6</sup> , R <sup>7</sup> , R <sup>8</sup> , R <sup>9</sup> , and R <sup>10</sup> is -L-M,
7		wherein
8		L is a linker independently selected from a bond, substituted or unsubstituted
9		alkylene, substituted or unsubstituted heteroalkylene, substituted or
10		unsubstituted cycloalkylene, substituted or unsubstituted
11		heterocycloalkylene, substituted or unsubstituted arylene, substituted or
12		unsubstituted heteroarylene, -O-, -NH-, -S-, and CH <sub>2</sub> -,
13		M is a modifying group, and
14		R <sup>11</sup> , R <sup>12</sup> , R <sup>13</sup> , R <sup>14</sup> , R <sup>15</sup> , and R <sup>16</sup> are independently selected from hydrogen,
15		substituted or unsubstituted alkyl, substituted or unsubstituted heteroalkyl,
16		and $-L^1-M^1$ , wherein
17		L <sup>1</sup> is a linker independently selected from a bond, substituted or
18		unsubstituted alkylene, substituted or unsubstituted heteroalkylene,

substituted or unsubstituted cycloalkylene, substituted or unsubstituted
heterocycloalkylene, substituted or unsubstituted arylene, substituted
or unsubstituted heteroarylene, -O-, -NH-, -S-, and CH<sub>2</sub>-, and
M<sup>1</sup> is modifying group.

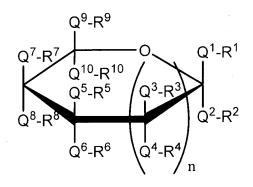
13. (Original) The method of claim 11, wherein said modified sugar has the formula

wherein,

W, X, Y, Z, and A are members independently selected from a bond, substituted or unsubstituted alkylene, substituted or unsubstituted or unsubstituted heteroalkylene, substituted or unsubstituted or unsubstituted heterocycloalkylene, substituted or unsubstituted arylene, substituted or unsubstituted heteroarylene, -O-, -N(R<sup>7</sup>)-, -S-, and -CH<sub>2</sub>-, wherein, R<sup>7</sup> is a member independently selected from hydrogen, substituted or unsubstituted alkyl, substituted or unsubstituted heteroalkyl, substituted or unsubstituted heterocycloalkyl, substituted or unsubstituted or unsubstituted heterocycloalkyl, substituted or unsubstituted or unsubstituted heteroaryl; and R<sup>1</sup>, R<sup>2</sup>, R<sup>3</sup>, R<sup>4</sup>, R<sup>5</sup> and R<sup>6</sup> are members independently selected from -OH, -

R<sup>1</sup>, R<sup>2</sup>, R<sup>3</sup>, R<sup>4</sup>, R<sup>5</sup> and R<sup>6</sup> are members independently selected from -OH, -NH<sub>2</sub>, -SH, hydrogen, substituted or unsubstituted alkyl, substituted or unsubstituted heteroalkyl, substituted or unsubstituted cycloalkyl, substituted or unsubstituted or unsubstituted aryl, substituted or unsubstituted heteroaryl, and a modifying group, wherein at least one or R<sup>1</sup>, R<sup>2</sup>, R<sup>3</sup>, R<sup>4</sup>, R<sup>5</sup> and R<sup>6</sup> is a modifying group.

14. (Original) The method of claim 4, wherein said modified nucleotide sugar has the formula



4 wherein,

n represents an integer from 0 to 1;

Q<sup>1</sup>, Q<sup>2</sup>, Q<sup>3</sup>, Q<sup>4</sup>, Q<sup>5</sup>, Q<sup>6</sup>, Q<sup>7</sup>, Q<sup>8</sup>, Q<sup>9</sup>, and Q<sup>10</sup> are members independently selected from a bond, substituted or unsubstituted alkylene, substituted or unsubstituted heteroalkylene, substituted or unsubstituted cycloalkylene, substituted or unsubstituted arylene, substituted or unsubstituted heterocycloalkylene, substituted or unsubstituted arylene, substituted or unsubstituted heteroarylene, -O-, -N(R<sup>1A</sup>)-, -S-, -C(O)-, and -CH<sub>2</sub>-, wherein

(I)

R<sup>1A</sup> is a member selected from hydrogen, substituted or unsubstituted alkyl, substituted or unsubstituted heteroalkyl, substituted or unsubstituted cycloalkyl, substituted or unsubstituted heterocycloalkyl, substituted or unsubstituted aryl, and substituted or unsubstituted heteroaryl; and

R<sup>1</sup>, R<sup>2</sup>, R<sup>3</sup>, R<sup>4</sup>, R<sup>5</sup>, R<sup>6</sup>, R<sup>7</sup>, R<sup>8</sup>, R<sup>9</sup>, and R<sup>10</sup> are members independently selected from -OPO<sub>3</sub>H<sub>2</sub>, -OH, -NH<sub>2</sub>, -SH, hydrogen, substituted or unsubstituted alkyl, substituted or unsubstituted heteroalkyl, substituted or unsubstituted cycloalkyl, substituted or unsubstituted heterocycloalkyl, substituted or unsubstituted aryl, substituted or unsubstituted heteroaryl, an activated leaving group, a nucleotidyl moiety, and a modifying group, wherein at least one of R<sup>1</sup>, R<sup>2</sup>, R<sup>3</sup>, R<sup>4</sup>, R<sup>5</sup>, R<sup>6</sup>, R<sup>7</sup>, R<sup>8</sup>, R<sup>9</sup>, and R<sup>10</sup> is a modifying group and a nucleotidyl moiety.

15. (Original) The method of claim 14, wherein said modified nucleotide sugar has the formula

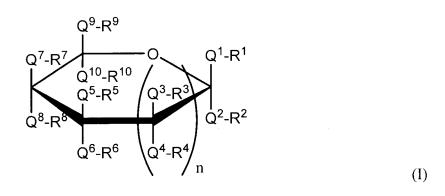
wherein,

W, X, Y, Z, and A are members independently selected from a bond, substituted or unsubstituted alkylene, substituted or unsubstituted or unsubstituted heteroalkylene, substituted or unsubstituted or unsubstituted or unsubstituted or unsubstituted or unsubstituted heteroarylene, -O-, -N(R<sup>7</sup>)-, -S-, and -CH<sub>2</sub>-, wherein, R<sup>7</sup> is a member independently selected from hydrogen, substituted or unsubstituted alkyl, substituted or unsubstituted heteroalkyl, substituted or unsubstituted heterocycloalkyl, substituted or unsubstituted heterocycloalkyl, substituted or unsubstituted or unsubstituted heteroaryl; and

R<sup>1</sup>, R<sup>2</sup>, R<sup>3</sup>, R<sup>4</sup>, and R<sup>5</sup> are independently selected from -OH, -NH<sub>2</sub>, -SH, hydrogen, substituted or unsubstituted alkyl, substituted or unsubstituted heteroalkyl, substituted or unsubstituted or unsubstituted or unsubstituted aryl, substituted or unsubstituted heteroaryl, and a modifying group, wherein at least one or R<sup>1</sup>, R<sup>2</sup>, R<sup>3</sup>, R<sup>4</sup>, and R<sup>5</sup> is a modifying group.

16. (Original) The method of claim 5, wherein said modified nucleotide sugar has the

## 2 formula



wherein, 4 5 n represents an integer from 0 to 1; Q<sup>1</sup>, Q<sup>2</sup>, Q<sup>3</sup>, Q<sup>4</sup>, Q<sup>5</sup>, Q<sup>6</sup>, Q<sup>7</sup>, Q<sup>8</sup>, Q<sup>9</sup>, and Q<sup>10</sup> are members independently selected 6 from a bond, substituted or unsubstituted alkylene, substituted or 7 unsubstituted heteroalkylene, substituted or unsubstituted cycloalkylene, 8 substituted or unsubstituted heterocycloalkylene, substituted or unsubstituted 9 arylene, substituted or unsubstituted heteroarylene, -O-, -N(R<sup>1A</sup>)-, -S-, -C(O)-, 10 and -CH<sub>2</sub>-, wherein 11 R<sup>1A</sup> is a member selected from hydrogen, substituted or unsubstituted alkyl, 12 substituted or unsubstituted heteroalkyl, substituted or unsubstituted 13 cycloalkyl, substituted or unsubstituted heterocycloalkyl, substituted or 14 unsubstituted aryl, and substituted or unsubstituted heteroaryl; and 15 R<sup>1</sup>, R<sup>2</sup>, R<sup>3</sup>, R<sup>4</sup>, R<sup>5</sup>, R<sup>6</sup>, R<sup>7</sup>, R<sup>8</sup>, R<sup>9</sup>, and R<sup>10</sup> are members independently selected 16 from -OPO<sub>3</sub>H<sub>2</sub>, -OH, -NH<sub>2</sub>, -SH, hydrogen, substituted or unsubstituted 17 alkyl, substituted or unsubstituted heteroalkyl, substituted or unsubstituted 18 cycloalkyl, substituted or unsubstituted heterocycloalkyl, substituted or 19 unsubstituted aryl, substituted or unsubstituted heteroaryl, an activated 20 leaving group, a nucleotidyl moiety, and a modifying group, wherein at least 21 one of R<sup>1</sup>, R<sup>2</sup>, R<sup>3</sup>, R<sup>4</sup>, R<sup>5</sup>, R<sup>6</sup>, R<sup>7</sup>, R<sup>8</sup>, R<sup>9</sup>, and R<sup>10</sup> is a modifying group and an 22 activated leaving group. 23 (Original) The method of claim 1, wherein said peptide is selected from the group 1 17. consisting of granulocyte colony stimulating factor, interferon-alpha, interferon-beta, Factor 2 VIIa, Factor IX, follicle stimulating hormone, erythropoietin, granulocyte macrophage colony 3 stimulating factor, interferon-gamma, alpha-1-protease inhibitor, glucocerebrosidase, tissue 4 plasminogen activator protein, interleukin-2, Factor VIII, chimeric tumor necrosis factor 5 receptor, urokinase, chimeric anti-glycoprotein IIb/IIIa antibody, chimeric anti-HER2 antibody, 6 chimeric anti-respiratory syncytial virus antibody, chimeric anti-CD20 antibody, DNase, 7 chimeric anti-tumor necrosis factor antibody, human insulin, hepatitis B sAg, interferon-omega, 8 alpha-galactosidase A, alpha-iduronidase, anti-thrombin III, human chorionic gonadotropin, and 9

human growth hormone.

10

1	<b>18</b> .	(Original) A cell comprising a peptide conjugate, said peptide conjugate comprising:
2		(i) a modifying group and a peptide, wherein said modifying group is linked to said
3		peptide via a glycosyl linking group interposed between and covalently linked to
4		both the peptide and said modifying group; and
5		(ii) said modifying group is a member independently selected from the group consisting
6		of a water-soluble polymer, a therapeutic moiety, a detectable label, and a
7		targeting moiety.
1	19.	(Original) The method of claim 18, wherein said glycosyl linking group is an intact
2	glyco	syl linking group.